SEQUENCE LISTING

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<120> Methods for Using Artificial Polynucleotides and Compositions thereof to Reduce Transgene Silencing

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<141> 2002-07-18

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ggattgaagc agcttggtgc agatgttgat tgtttccttg gcactgactg cccacctgtt 660

cgtgtcaatg gaatcggagg gctacctggt ggcaaggtca agctgtctgg ctccatcagc 720

agtcagtact tgagtgcctt gctgatggct gctcctttgg ctcttgggga tgtggagatt 780

gaaatcattg ataaattaat ctccattccg tacgtcgaaa tgacattgag attgatggag 840

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gtcaccatcc gggaccctgg gtgcacccgg aagaccttcc ccgactactt cgatgtgctg 1500

agcactttcg tcaagaatta a 1521

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<212> DNA

<213> Zea mays

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cttagagcgc ctggtagaag ggttatcgca gcgccaccgg cggctgccgc tgcggcagcg 180

gtgcaggccg gcgcggaaga gatcgtccta cagcctatca aggaaatctc tggtacggta 240

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gggaccacgg tcgtagataa tctgctcaac agcgaagacg tgcactatat gttgggtgcc 360

ctgaggacgc taggtctgtc agtggaagcc gataaggccg ccaagcgcgc tgtcgtcgtt 420

ggctgcggcg gtaagttccc cgtggaggac gcgaaagaag aggtgcagtt atttcttggg 480

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gagggcccgg attactgcat aatcacacct ccagagaagt tgaacgtcac cgctatcgac 1380

acatacgacg atcaccggat ggcaatggcc tttagcttgg cagcgtgcgc cgaagtacct 1440

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tcaaccttcg tgaagaactg a 1521

<210> 11
<211> 76
<212> PRT
<213> Arabidopsis thaliana

<400> 11

Met Ala Gln Val Ser Arg Ile Cys Asn Gly Val Gln Asn Pro Ser Leu
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Ile Ser Asn Leu Ser Lys Ser Ser Gln Arg Lys Ser Pro Leu Ser Val
20 25 30

Ser Leu Lys Thr Gln Gln His Pro Arg Ala Tyr Pro Ile Ser Ser Ser
35 40 45

Trp Gly Leu Lys Lys Ser Gly Met Thr Leu Ile Gly Ser Glu Leu Arg
50 55 60

Pro Leu Lys Val Met Ser Ser Val Ser Thr Ala Cys
65 70 75

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<211> 228
<212> DNA
<213> Arabidopsis thaliana

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cgagcttatc cgatttcgtc gtcgtgggga ttgaagaaga gtgggatgac gttaattggc 180

tctgagcttc gtcctcttaa ggtcatgtct tctgtttcca cggcgtgc 228

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<213> Arabidopsis thaliana

<400> 13
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agggcttatc ccatctcaag ctcctggggt ctaaagaaaa gtggaatgac actgatcggt 180

agcgaactac gaccgctgaa agtcatgtcc tcagtcagca ctgcgtgc 228

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<213> Arabidopsis thaliana

<400> 14
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agggcctacc ctatcagctc atcctggggc ctcaagaaga gtggcatgac gctgatcggc 180

agcgagctgc ggccactcaa ggtgatgtcc tcggtctcaa cggcgtgc 228

<210> 15
<211> 455
<212> PRT
<213> Agrobacterium tumefaciens

<400> 15

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Gly Leu Ser Gly Thr Val Arg Ile Pro Gly Asp Lys Ser Ile Ser His
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Arg Ser Phe Met Phe Gly Gly Leu Ala Ser Gly Glu Thr Arg Ile Thr
35 40 45

Gly Leu Leu Glu Gly Glu Asp Val Ile Asn Thr Gly Lys Ala Met Gln
50 55 60

Ala Met Gly Ala Arg Ile Arg Lys Glu Gly Asp Thr Trp Ile Ile Asp
65 70 75 80

Gly Val Gly Asn Gly Gly Leu Leu Ala Pro Glu Ala Pro Leu Asp Phe
85 90 95

Gly Asn Ala Ala Thr Gly Cys Arg Leu Thr Met Gly Leu Val Gly Val
100 105 110

Tyr Asp Phe Asp Ser Thr Phe Ile Gly Asp Ala Ser Leu Thr Lys Arg
115 120 125

Pro Met Gly Arg Val Leu Asn Pro Leu Arg Glu Met Gly Val Gln Val
130 135 140

Lys Ser Glu Asp Gly Asp Arg Leu Pro Val Thr Leu Arg Gly Pro Lys
145 150 155 160

Thr Pro Thr Pro Ile Thr Tyr Arg Val Pro Met Ala Ser Ala Gln Val
165 170 175

Lys Ser Ala Val Leu Leu Ala Gly Leu Asn Thr Pro Gly Ile Thr Thr
180 185 190

Val Ile Glu Pro Ile Met Thr Arg Asp His Thr Glu Lys Met Leu Gln
195 200 205

Gly Phe Gly Ala Asn Leu Thr Val Glu Thr Asp Ala Asp Gly Val Arg
210 215 220

Thr Ile Arg Leu Glu Gly Arg Gly Lys Leu Thr Gly Gln Val Ile Asp
225 230 235 240

Val Pro Gly Asp Pro Ser Ser Thr Ala Phe Pro Leu Val Ala Ala Leu
245 250 255

Leu Val Pro Gly Ser Asp Val Thr Ile Leu Asn Val Leu Met Asn Pro
260 265 270

Thr Arg Thr Gly Leu Ile Leu Thr Leu Gln Glu Met Gly Ala Asp Ile
275 280 285

Glu Val Ile Asn Pro Arg Leu Ala Gly Gly Glu Asp Val Ala Asp Leu
290 295 300

Arg Val Arg Ser Ser Thr Leu Lys Gly Val Thr Val Pro Glu Asp Arg
305 310 315 320

Ala Pro Ser Met Ile Asp Glu Tyr Pro Ile Leu Ala Val Ala Ala Ala
325 330 335

Phe Ala Glu Gly Ala Thr Val Met Asn Gly Leu Glu Glu Leu Arg Val
340 345 350

Lys Glu Ser Asp Arg Leu Ser Ala Val Ala Asn Gly Leu Lys Leu Asn
355 360 365

Gly Val Asp Cys Asp Glu Gly Glu Thr Ser Leu Val Val Arg Gly Arg
370 375 380

Pro Asp Gly Lys Gly Leu Gly Asn Ala Ser Gly Ala Ala Val Ala Thr
385 390 395 400

His Leu Asp His Arg Ile Ala Met Ser Phe Leu Val Met Gly Leu Val
405 410 415

Ser Glu Asn Pro Val Thr Val Asp Asp Ala Thr Met Ile Ala Thr Ser
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Phe Pro Glu Phe Met Asp Leu Met Ala Gly Leu Gly Ala Lys Ile Glu

435 440 445

Leu Ser Asp Thr Lys Ala Ala
450 455

<210> 16
<211> 1368
<212> DNA
<213> Agrobacterium tumefaciens

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accgtccgca ttcccggcga caagtcgatc tcccaccggt ccttcatgtt cggcggtctc 120

gcgagcggtg aaacgcgcat caccggcctt ctggaaggcg aggacgtcat caatacgggc 180

aaggccatgc aggcgatggg cgcccgcatc cgtaaggaag gcgacacctg gatcatcgat 240

ggcgtcggca atggcggcct cctggcgcct gaggcgccgc tcgatttcgg caatgccgcc 300

acgggctgcc gcctgacgat gggcctcgtc ggggtctacg atttcgacag caccttcatc 360

ggcgacgcct cgctcacaaa gcgcccgatg ggccgcgtgt tgaacccgct gcgcgaaatg 420

ggcgtgcagg tgaaatcgga agacggtgac cgtcttcccg ttaccttgcg cgggccgaag 480

acgccgacgc cgatcaccta ccgcgtgccg atggcctccg cacaggtgaa gtccgccgtg 540

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gatcatacgg aaaagatgct gcagggcttt ggcgccaacc ttaccgtcga gacggatgcg 660

gacggcgtgc gcaccatccg cctggaaggc cgcggcaagc tcaccggcca agtcatcgac 720

gtgccgggcg acccgtcctc gacggccttc ccgctggttg cggccctgct tgttccgggc 780

tccgacgtca ccatcctcaa cgtgctgatg aaccccaccc gcaccggcct catcctgacg 840

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gtcacggtgg acgatgccac gatgatcgcc acgagcttcc cggagttcat ggacctgatg 1320

gccgggctgg gcgcgaagat cgaactctcc gatacgaagg ctgcctga 1368

<210> 17

<211> 1368
<212> DNA
<213> Agrobacterium tumefaciens

<400> 17
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accgtgcgca tccctggcga taaaagtatt tcacacagga gcttcatgtt cggaggactt 120

gctagtggag agacgagaat cactggtttg cttgagggcg aagatgttat caacaccggt 180

aaggcgatgc aagcaatggg tgccagaatc cgaaaagagg gcgatacgtg gatcatcgac 240

ggtgttggta acggaggatt gctcgctccc gaagcgccac ttgactttgg gaacgcagct 300

acggggtgcc gtcttactat gggactggta ggcgtgtatg actttgactc taccttcatc 360

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acccccacgc caatcacgta cagggttccg atggcgtcag cacaggtcaa gtcagcggta 540

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<210> 18
<211> 1368
<212> DNA
<213> Agrobacterium tumefaciens

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gccagcggtg agacgcgcat cacgggcctg cttgaaggtg aggacgtgat caataccggg 180

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gcgggcctcg gggccaagat cgaactgtct gacacgaagg ccgcttga 1368

<210> 19
<211> 183
<212> PRT
<213> Streptomyces hygroscopicus

<400> 19

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Asp Met Pro Ala Val Cys Thr Ile Val Asn His Tyr Ile Glu Thr Ser
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Thr Val Asn Phe Arg Thr Glu Pro Gln Glu Pro Gln Asp Trp Thr Asp
35 40 45

Asp Leu Val Arg Leu Arg Glu Arg Tyr Pro Trp Leu Val Ala Glu Val
50 55 60

Asp Gly Glu Val Ala Gly Ile Ala Tyr Ala Gly Pro Trp Lys Ala Arg
65 70 75 80

Asn Ala Tyr Asp Trp Thr Ala Glu Ser Thr Val Tyr Val Ser Pro Arg
85 90 95

His Gln Arg Thr Gly Leu Gly Ser Thr Leu Tyr Thr His Leu Leu Lys
100 105 110

Ser Leu Glu Ala Gln Gly Phe Lys Ser Val Val Ala Val Ile Gly Leu
115 120 125

Pro Asn Asp Pro Ser Val Arg Met His Glu Ala Leu Gly Tyr Ala Pro
130 135 140

Arg Gly Met Leu Arg Ala Ala Gly Phe Lys His Gly Asn Trp His Asp
145 150 155 160

Val Gly Phe Trp Gln Leu Asp Phe Ser Leu Pro Val Pro Pro Arg Pro
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Val Leu Pro Val Thr Glu Ile
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<210> 20
<211> 552
<212> DNA
<213> Streptomyces hygroscopicus

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<210> 21
<211> 552
<212> DNA
<213> Streptomyces hygroscopicus

<400> 21
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<210> 22
<211> 552
<212> DNA
<213> Streptomyces hygroscopicus

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<210> 23
<211> 1368
<212> DNA

<213> Agrobacterium tumefaciens

<400> 23
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<210> 24
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<212> DNA
<213> Artificial sequence

<220>
<223> DNA primer molecule

<400> 24
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<210> 25
<211> 16
<212> DNA
<213> Artificial sequence

<220>
<223> DNA primer molecule

<400> 25
gcctttgagt gtacta 16

<210> 26
<211> 16
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<220>
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